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Links

☐ 1: NM 075774. Caenorhabditis el...[gi:25150522] LOCUS mRNA INV 12-JUL-2003 NM 075774 1473 bp linear DEFINITION Caenorhabditis elegans Suppressor/Enhancer of Lin-12 SEL-12, SUppressor of Multi-vulva phenotype SUM-1, presenilin, membrane protein facilitator of Notch receptors signaling (50.0 kD) (sel-12) complete mRNA. ACCESSION NM 075774 VERSION NM 075774.2 GI:25150522 **KEYWORDS** SOURCE Caenorhabditis elegans (worm) ORGANISM Caenorhabditis elegans Eukaryota ; Metazoa ; Nematoda ; Chromadorea ; Rhabditida ; Rhabditoidea ; Rhabditidae ; Peloderinae ; Caenorhabditis. REFERENCE (bases 1 to 1473) AUTHORS Lakowski, B., Eimer, S., Gobel, C., Bottcher, A., Wagler, B. and Baumeister, R. TITLE Two suppressors of sel-12 encode C2H2 zinc-finger proteins that regulate presenilin transcription in Caenorhabditis elegans JOURNAL Development 130 (10), 2117-2128 (2003) MEDLINE 22554535 PUBMED 12668626 REFERENCE (bases 1 to 1473) AUTHORS Kitagawa, N., Shimohama, S., Oeda, T., Uemura, K., Kohno, R., Kuzuya, A., Shibasaki, H. and Ishii, N. TITLE The role of the presenilin-1 homologue gene sel-12 of Caenorhabditis elegans in apoptotic activities JOURNAL J. Biol. Chem. 278 (14), 12130-12134 (2003) 22552452 MEDLINE PUBMED 12556527 REFERENCE (bases 1 to 1473) Eimer, S., Donhauser, R. and Baumeister, R. **AUTHORS** The Caenorhabditis elegans presenilin sel-12 is required for TITLE mesodermal patterning and muscle function JOURNAL Dev. Biol. 251 (1), 178-192 (2002) MEDLINE 22301592 PUBMED 12413907 REFERENCE (bases 1 to 1473) Eimer, S., Lakowski, B., Donhauser, R. and Baumeister, R. AUTHORS TITLE Loss of spr-5 bypasses the requirement for the C.elegans presenilin sel-12 by derepressing hop-1 **JOURNAL** EMBO J. 21 (21), 5787-5796 (2002) MEDLINE 22299931 PUBMED 12411496 REFERENCE (bases 1 to 1473)

AUTHORS Li,J., Pauley,A.M., Myers,R.L., Shuang,R., Brashler,J.R., Yan,R., Buhl,A.E., Ruble,C. and Gurney,M.E.

SEL-10 interacts with presentilin 1, facilitates its ubiquitination, and alters A-beta peptide production

JOURNAL J. Neurochem. 82 (6), 1540-1548 (2002)

MEDLINE <u>22242246</u> PUBMED <u>12354302</u>

TITLE

REFERENCE 6 (bases 1 to 1473)

AUTHORS Francis, R., McGrath, G., Zhang, J., Ruddy, D.A., Sym, M., Apfeld, J., Nicoll, M., Maxwell, M., Hai, B., Ellis, M.C., Parks, A.L., Xu, W., Li, J., Gurney, M., Myers, R.L., Himes, C.S., Hiebsch, R., Ruble, C.,

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            aph-1 and pen-2 are required for Notch pathway signaling,
  TITLE
            gamma-secretase cleavage of betaAPP, and presenilin protein
            accumulation
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  JOURNAL
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            Levitan, D., Yu, G., St George Hyslop, P. and Goutte, C.
 AUTHORS
            APH-2/nicastrin functions in LIN-12/Notch signaling in the
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            Caenorhabditis elegans somatic gonad
            Dev. Biol. 240 (2), 654-661 (2001)
  JOURNAL
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            Maruyama, S., Hatakeyama, S., Nakayama, K., Ishida, N., Kawakami, K. and
 AUTHORS
            Nakayama, K.
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            of Caenorhabditis elegans SEL-10
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  JOURNAL
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REFERENCE
            Cinar, H.N., Sweet, K.L., Hosemann, K.E., Earley, K. and Newman, A.P.
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            10 (bases 1 to 1473)
  AUTHORS
            Okochi, M., Eimer, S., Bottcher, A., Baumeister, R., Romig, H.,
            Walter, J., Capell, A., Steiner, H. and Haass, C.
            A loss of function mutant of the presenilin homologue SEL-12
  TITLE
            undergoes aberrant endoproteolysis in Caenorhabditis elegans and
            increases abeta 42 generation in human cells
  JOURNAL
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REFERENCE
            11 (bases 1 to 1473)
  AUTHORS
            Wen, C., Levitan, D., Li, X. and Greenwald, I.
  TITLE
            spr-2, a suppressor of the egg-laying defect caused by loss of
            sel-12 presenilin in Caenorhabditis elegans, is a member of the SET
            protein subfamily
  JOURNAL
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 AUTHORS
            Zhang, D.M., Levitan, D., Yu, G., Nishimura, M., Chen, F., Tandon, A.,
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            Holmes, E., Milman, P., Sato, C., Zhang, L. and St George-Hyslop, P.
  TITLE
            Mutation of the conserved N-terminal cysteine (Cys92) of human
            presenilin 1 causes increased A beta42 secretion in mammalian cells
            but impaired Notch/lin-12 signalling in C. elegans
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            Yu,G., Nishimura,M., Arawaka,S., Levitan,D., Zhang,L., Tandon,A.,
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            Aebersold, R., Farrer, L.S., Sorbi, S., Bruni, A., Fraser, P. and St
            George-Hyslop, P.
  TITLE
            Nicastrin modulates presenilin-mediated notch/glp-1 signal
            transduction and betaAPP processing
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            Wittenburg, N., Eimer, S., Lakowski, B., Rohrig, S., Rudolph, C. and
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            Baumeister, R.
  TITLE
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            neurons in C. elegans
  JOURNAL
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REFERENCE
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  AUTHORS
            Kurt, H., Grim, M.G., Baumeister, R. and Loetscher, H.
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  TITLE
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            16 (bases 1 to 1473)
  AUTHORS
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            Shen, J., Lu, F.M., Lux, S.E., Tonegawa, S. and Hyman, B.T.
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            primary mammalian neurons
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            Brain Res. Mol. Brain Res. 69 (2), 273-280 (1999)
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REFERENCE
            17 (bases 1 to 1473)
            Ray, W.J., Yao, M., Nowotny, P., Mumm, J., Zhang, W., Wu, J.Y., Kopan, R.
  AUTHORS
            and Goate, A.M.
            Evidence for a physical interaction between presentlin and Notch
  TITLE
            Proc. Natl. Acad. Sci. U.S.A. 96 (6), 3263-3268 (1999)
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REFERENCE
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  AUTHORS
            Westlund, B., Parry, D., Clover, R., Basson, M. and Johnson, C.D.
  TITLE
            Reverse genetic analysis of Caenorhabditis elegans presenilins
            reveals redundant but unequal roles for sel-12 and hop-1 in
            Notch-pathway signaling
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  AUTHORS
            Hong, C.S., Caromile, L., Nomata, Y., Mori, H., Bredesen, D.E. and
            Koo, E.H.
  TITLE
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            differentiation in vitro
            J. Neurosci. 19 (2), 637-643 (1999)
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            20 (bases 1 to 1473)
  AUTHORS
            Wu, G., Hubbard, E.J., Kitajewski, J.K. and Greenwald, I.
  TITLE
            Evidence for functional and physical association between
            Caenorhabditis elegans SEL-10, a Cdc4p-related protein, and SEL-12
            presenilin
  JOURNAL
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REFERENCE
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  AUTHORS
            Levitan, D. and Greenwald, I.
  TITLE
            Effects of SEL-12 presentlin on LIN-12 localization and function in
            Caenorhabditis elegans
  JOURNAL
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            22 (bases 1 to 1473)
            Berezovska, O., Xia, M.Q. and Hyman, B.T.
  AUTHORS
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            presenilin-1, and is altered in Alzheimer disease
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REFERENCE
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 AUTHORS
            Li, X. and Greenwald, I:
            Additional evidence for an eight-transmembrane-domain topology for
  TITLE
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  JOURNAL
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  AUTHORS
            Davis, J.A., Naruse, S., Chen, H., Eckman, C., Younkin, S., Price, D.L.,
            Borchelt, D.R., Sisodia, S.S. and Wong, P.C.
            An Alzheimer's disease-linked PS1 variant rescues the developmental
  TITLE
            abnormalities of PS1-deficient embryos
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REFERENCE
            Zhang, W., Han, S.W., McKeel, D.W., Goate, A. and Wu, J.Y.
 AUTHORS
            Interaction of presenilins with the filamin family of actin-binding
  TITLE .
            J. Neurosci. 18 (3), 914-922 (1998)
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            26 (bases 1 to 1473)
            Mattson, M.P., Guo, Q., Furukawa, K. and Pedersen, W.A.
  AUTHORS
  TITLE
            Presenilins, the endoplasmic reticulum, and neuronal apoptosis in
            Alzheimer's disease
            J. Neurochem. 70 (1), 1-14 (1998)
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            27 (bases 1 to 1473)
REFERENCE
  AUTHORS
            Mattson, M.P. and Guo, Q.
  TITLE
            Cell and molecular neurobiology of presenilins: a role for the
            endoplasmic reticulum in the pathogenesis of Alzheimer's disease?
  JOURNAL
            J. Neurosci. Res. 50 (4), 505-513 (1997)
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REFERENCE
            28 (bases 1 to 1473)
  AUTHORS
            Li, X. and Greenwald, I.
  TITLE
            HOP-1, a Caenorhabditis elegans presenilin, appears to be
            functionally redundant with SEL-12 presentlin and to facilitate
            LIN-12 and GLP-1 signaling
  JOURNAL
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  AUTHORS
            Hutton, M. and Hardy, J.
  TITLE
            The presentlins and Alzheimer's disease
            Hum. Mol. Genet. 6 (10), 1639-1646 (1997)
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            30 (bases 1 to 1473)
  AUTHORS
            Wong, P.C., Zheng, H., Chen, H., Becher, M.W., Sirinathsinghji, D.J.,
            Trumbauer, M.E., Chen, H.Y., Price, D.L., Van der Ploeg, L.H. and
            Sisodia, S.S.
  TITLE
            Presenilin 1 is required for Notch1 and DII1 expression in the
            paraxial mesoderm
            Nature 387 (6630), 288-292 (1997)
  JOURNAL
            97297761
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            31 (bases 1 to 1473)
  AUTHORS
            Baumeister, R., Leimer, U., Zweckbronner, I., Jakubek, C., Grunberg, J.
            and Haass, C.
  TITLE
            Human presenilin-1, but not familial Alzheimer's disease (FAD)
            mutants, facilitate Caenorhabditis elegans Notch signalling
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independently of proteolytic processing Genes Funct. 1 (2), 149-159 (1997) JOURNAL MEDLINE 98343909 PUBMED 9680315 REFERENCE 32 (bases 1 to 1473) AUTHORS Hong, C.S. and Koo, E.H. TITLE Isolation and characterization of Drosophila presentlin homolog JOURNAL Neuroreport 8 (3), 665-668 (1997) 97260623 MEDLINE PUBMED 9106743 33 (bases 1 to 1473) REFERENCE AUTHORS Berezovska, O., Xia, M.Q., Page, K., Wasco, W., Tanzi, R.E. and Hyman, B.T. TITLE Developmental regulation of presentlin mRNA expression parallels notch expression JOURNAL J. Neuropathol. Exp. Neurol. 56 (1), 40-44 (1997) MEDLINE 97144360 PUBMED 8990127 REFERENCE 34 (bases 1 to 1473) Levitan, D., Doyle, T.G., Brousseau, D., Lee, M.K., Thinakaran, G., AUTHORS Slunt, H.H., Sisodia, S.S. and Greenwald, I. TITLE Assessment of normal and mutant human presentlin function in Caenorhabditis elegans JOURNAL Proc. Natl. Acad. Sci. U.S.A. 93 (25), 14940-14944 (1996) MEDLINE 97121494 PUBMED 8962160 REFERENCE 35 (bases 1 to 1473) AUTHORS Li, X. and Greenwald, I. TITLE Membrane topology of the C. elegans SEL-12 presenilin JOURNAL Neuron 17 (5), 1015-1021 (1996) MEDLINE 97092712 PUBMED 8938132 REFERENCE 36 (bases 1 to 1473) AUTHORS Levitan, D. and Greenwald, I. TITLE Facilitation of lin-12-mediated signalling by sel-12, a Caenorhabditis elegans S182 Alzheimer's disease gene JOURNAL Nature 377 (6547), 351-354 (1995) MEDLINE 96032531 PUBMED 7566091 COMMENT REVIEWED REFSEQ: This record has been curated by NCBI staff. The reference sequence was derived from U35660 and AV179958.1. On Nov 21, 2002 this sequence version replaced gi:17569442. Summary: This gene sel-12, also known as sum-1, F35H12.3, XB535 or YK4554, maps at (X; -19.01). Its phenotype is suppressor/enhancer of lin-12, suppressor of multi-vulva phenotype, facilitator of notch-type receptors signaling. It encodes a presenilin, membrane protein facilitator of Notch receptors signaling. From Pfam homology, the product would be involved in intracellular signaling cascade and would localize in membrane.

According to the Worm Transcriptome Project, it is well expressed in L3, L4, adult and culminating in embryos [Kohara cDNAs]. Its sequence is defined by 11 cDNA clones.

### Phenotype

[from C. elegans II book] Allele ar131: (previously known as sum-1) recessive suppressor of multivulva phenotype of lin-12 hypermorph n950; impenetrant egg laying defective in lin-12 (+) background. Three other alleles: ar133, ar171 (100% egg laying defective, ar171/Df similar, W225opal). Cloned: encodes predicted 467 aa protein, 9 transmembrane domains; related to human presenilin genes (S182) and to SPE-4. [Levitan and Greenwald 1995; Iva Greenwald]. Allele ar131, ar40.

[Levitan D] suppressor of multivulva phenotype. Selected strains available from the CGC.

GS883 dpy-5(e61) sel(ar40)I; unc-32(e189) lin-12(n676n930)III

[Greenwald IS] DpyUnc. ar40 is a semi-dominant suppressor. At 25C ar40 suppresses the Egl phenotype of ne676n930. At 15C a high percentage of hermaphrodites have a 0 AC-Egl phenotype. ar40 suppresses proximal mitosis. ar40 does not suppress vulval lineage defects.

AN87 sel-12(tyll) X [Anna Newman, Nese Cinar, EMS] Egl. Premature stop codon.

#### RNA interference results:

[J.Ahringer 2003] No obvious phenotype (by feeding genomic PCR product JA:F35H12.3). Warning: this double stranded RNA may also interfere with gene XB537.

#### Function

Protein properties: [GB:AF171064] function: facilitator of Notch receptors signaling.

membrane protein similar to Homo sapiens PS1 and PS2.

[WormBase] The sel-12 gene encodes a ortholog of human PS1, which when mutated leads to type 3 Alzheimer disease (OMIM:104311); it is also homologous to PS2, which when mutated leads type 4 Alzheimer disease (OMIM:600759).

#### Expression

The expression profile for the gene, derived from the proportion of animals at each stage in each Kohara library is: embryos 76%, L1 or L2 larvae 1%, L3 to adult 22%.

In situ hybridisation pictures to all stages of development are available from Kohara NextDB.

For a detailed expression pattern description, see Wormbase Expr1288, Expr1609.

#### Interactions

eukaryota.

This gene interacts with:

gene spr-1: spr-1 loss of function suppresses Egl of sel-12. protein LIN-12. protein SEL-10CO.

This complete mRNA is 1473 bp long. Its sequence exactly matches the genome. The premessenger has 7 exons. It covers 2.42 kb on the WS97 genome. It is transpliced to SL1. The protein (444 aa, 50.0 kDa, pI 6.7) contains one Presentlin motif. It also contains at least 8 transmembrane domain, a prenylation domain, an ER membrane domain [Psort2]. It is predicted to localise in the plasma membrane [Psort2]. Taxblast (threshold 10^-3) tracks ancestors down to

COMPLETENESS: full length.

## FEATURES

source

Location/Qualifiers

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yk674e3, yk499e3, yk400e8, yk600e12, yk216e1, yk231a7, yk573h4, yk452b9. for edited clone sequences see www.wormgenes.org"

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containing 15-30% males: yk231a7; gb: AF171064, U35660"
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                 /gene="sel-12"
                 /locus_tag="XB535"
                 /note="Region: [PSORT] ER membrane domain: KCLL"
misc feature
                 1321..1332
                 /gene="sel-12"
                 /locus tag="XB535"
                 /note="Region: [PSORT] prenylation domain: CLLY"
                 1..54
exon
                 /gene="sel-12"
                 /locus tag="XB535"
                 /note="Exon 1 length 54 bp"
misc feature
                bond (54,55)
                 /gene="sel-12"
                 /locus tag="XB535"
                 /note="Intron length 70 bp, type gt_ag"
                55..245
exon
                 /gene="sel-12"
                 /locus tag="XB535"
                 /note="Exon 2 length 191 bp"
misc feature
                bond (245, 246)
                 /gene="sel-12"
                 /locus_tag="XB535"
                 /note="Intron length 304 bp, type gt ag"
                 246..455
exon
                 /gene="sel-12"
                 /locus tag="XB535"
                 /note="Exon 3 length 210 bp"
misc feature
                bond (455, 456)
                 /qene="sel-12"
                 /locus tag="XB535"
                 /note="Intron length 45 bp, type gt_ag"
                456..775
exon
                 /gene="sel-12"
                 /locus_tag="XB535"
                 /note="Exon 4 length 320 bp"
misc feature
                bond(775,776)
                /gene="sel-12"
                /locus_tag="XB535"
                /note="Intron length 61 bp, type gt_ag"
                776..859
exon
                /gene="sel-12"
                /locus tag="XB535"
                /note="Exon 5 length 84 bp"
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```
bond (859, 860)
     misc reature
                     /gene="sel-12"
                     /locus tag="XB535"
                     /note="Intron length 49 bp, type gt_ag"
                     860..1066
     exon
                     /gene="sel-12"
                     /locus tag="XB535"
                     /note="Exon 6 length 207 bp"
                     bond (1066, 1067)
     misc feature
                     /gene="sel-12"
                     /locus_tag="XB535"
                     /note="Intron length 422 bp, type gt_ag"
                     1067..1473
     exon
                     /gene="sel-12"
                     /locus tag="XB535"
                     /note="Exon 7 length 407 bp"
     3'UTR
                     1336..1473
                     /gene="sel-12"
                     /locus tag="XB535"
                     /note="The 3' UTR contains 138 bp followed by the polyA.
                     The standard AATAAA polyadenylation signal does not occur,
                     but the variant ATTAAA is seen about 15 bp before the
                     polyA."
                     /evidence=experimental
     polyA signal
                     1459..1464
                     /gene="sel-12"
                     /locus tag="XB535"
                     /note="variant attaaa"
                     1473
     polyA site
                     /qene="sel-12"
                     /locus tag="XB535"
                     /note="PolyA visible in U35660, yk452b9"
                     /evidence=experimental
BASE COUNT
                381 a
                         313 c
                                  312 q
                                           467 t
ORIGIN
        1 atgcetteca caaggagaca acaggaggge ggaggtgcag atgcggaaac acataccgtt
       61 tacggtacaa atctgataac aaatcggaat agccaagaag acgaaaatgt tgtggaagaa
      121 geggagetga aataeggage ateteaegtt atteatetat ttgtgeeggt gteaetatge
      181 atggctctgg ttgtttttac gatgaacacg attacgtttt atagtcaaaa caatggaagg
      241 catttactat acacteettt tgteegggaa acagacagta tegttgagaa gggattgatg
      301 tcacttggaa atgetetegt catgttgtge gtggtegtte tgatgacagt tetgetgatt
      361 gttttctata aatacaagit ttataagctt attcatggat ggcttattgt cagcagtttt
      421 cttcttcttt tcctattcac tacaatctat gtgcaagaag ttctgaaaag tttcgatgtg
      481 teteccageg caetattggt tttgtttgga etgggtaaet atggagttet eggaatgatg
      541 tgtatacatt ggaaaggtee attgegtetg caacagttet acettattac aatgtetgea 🤙
      601 ctaatggctc tggtctttat caagtaccta ccagaatgga ctgtgtggtt tgtgctgttt
      661 gttatctcgg tttgggatct ggttgccgtg ctcacaccaa aaggaccatt gagatatttg
      721 gtggaaactg cacaggagag aaacgagcca attttcccgg cgctgattta ttcgtctgga
      781 gtcatctatc cctacgttct tgttactgca gttgaaaaca cgacagaccc ccqtqaaccq
      841 acgtcgtcag actcaaatac ttctacagct tttcctggag aggcgagttg ttcatctgaa
      901 acgccaaaac ggccaaaagt gaaacgaatt cctcaaaaag tgcaaatcga atcgaatact
      961 acagetteaa egacacaaaa etetggagta agggtggaac gggagetage tgetgagaga
     1021 ccaactgtac aagacgccaa ttttcacagg cacgaagagg aagagagagg tgtgaaactt
     1081 ggtctgggcg acttcatttt ctactctgtt ctcctcggca aggcttcatc gtactttgac
     1141 tggaacacga ctatcgcttg ttatgtggcc attcttatcg gtctctgctt cactcttgtc
     1201 ctgctcgccg tcttcaaacg agcactcccg gctctgccaa tttccatttt ctccggactc
     1261 attitttact titgtacecg ciggateate accepatitg tiacacaagt cicteaaaag
     1321 tgtttattat attaattete tgtttttgee atttetttge atcateaact tttegattat
     1381 atcttgagcg atctcaaagc tttattttac atacctattt atttttgaac tttgtcattt
     1441 aagttatata aataatttat taaacgtttc tgc
```

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• Nature. 1995 Jun 29;375(6534):734.

Cloning of a gene bearing missense mutations in early-onset familial Alzheimer's disease.

Sherrington R, Rogaev EI, Liang Y, Rogaeva EA, Levesque G, Ikeda M, Chi H, Lin C, Li G, Holman K, et al.

Department of Medicine (Neurology), University of Toronto, Ontario, Canada.

Some cases of Alzheimer's disease are inherited as an autosomal dominant trait. Genetic linkage studies have mapped a locus (AD3) associated with susceptibility to a very aggressive form of Alzheimer's disease to chromosome 14q24.3. We have defined a minimal cosegregating region containing the AD3 gene, and isolated at least 19 different transcripts encoded within this region. One of these transcripts (S182) corresponds to a novel gene whose product is predicted to contain multiple transmembrane domains and resembles an integral membrane protein. Five different missense mutations have been found that cosegregate with early-onset familial Alzheimer's disease. Because these changes occurred in conserved domains of this gene, and are not present in normal controls, they are likely to be causative of AD3.

#### MeSH Terms:

- Alzheimer Disease/genetics\*
- Amino Acid Sequence
- Animal
- Base Sequence
- Chromosome Mapping
- Chromosomes, Human, Pair 14\*
- Cloning, Molecular\*
- Female
- Human
- Male
- Membrane Proteins/chemistry
- Membrane Proteins/genetics\*
- Mice
- Molecular Sequence Data
- Mutation\*
- Open Reading Frames
- Pedigree
- Protein Structure, Secondary
- Support, Non-U.S. Gov't
- Transcription, Genetic

# Gene Symbols:

- AD3
- S182

### Substances:

- Membrane Proteins
- S182 protein

# Secondary Source ID:

- GENBANK/L40391
- GENBANK/L40392
- GENBANK/L40393
- GENBANK/L40394
- GENBANK/L40395
- GENBANK/L40396
- GENBANK/L40397
- GENBANK/L40398
- GENBANK/L40399
- GENBANK/L40400
- GENBANK/L40401
- GENBANK/L40402
- GENBANK/L40403
- GENBANK/L42110
- GENBANK/L42110
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- GENBANK/L76517
- GENBANK/L76518
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